## IN THE CLAIMS

This listing of claims replaces all prior versions, and listings, in this application.

Claims 1-12 (canceled)

- 13. (withdrawn/currently amended) A method for transferring N-acetyl-D-glucosamine from a donor substrate to an acceptor substrate through  $\beta$ 1,3-linkage, wherein " $\beta$ " represents an anomer assuming a cis configuration, of anomers of glycosidic linkage at position 1 of the sugar ring, the method comprising reacting the donor substrate and the acceptor substrate with a  $\beta$ 1,3-N-acetyl-D-glucosaminyltransferase protein, wherein the protein comprises the following-amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 16. [[:]]
- (A) SEQ ID NO: 2, SEQ ID NO: 16, or SEQ ID NO: 17; or
- (B) SEQ ID NO: 2, SEQ ID NO: 16, or SEQ ID NO: 17 in which one or to 20 amine acid(s) is(are) substituted, deleted, or inserted.
- 14. (currently amended) An isolated <u>β1,3-N-acetyl-D- glucosaminyltransferase protein</u> comprising the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 16 glycosyltransferase protein, wherein the protein has at least one of the following properties (a) to (c):
- (a) acceptor substrate specificity: the protein transfers GlcNAc from UDP-GlcNAc to an acceptor substrate having an oligosaccharide residue in a quadruple-stranded form at the nonreducing end of an N-linked oligosaccharide through a β1,3 glycosidic linkage and synthesizes an oligosaccharide has a significant transferring activity for at least Bz-β-lactoside and/or Galβ1-4GlcNAc groups, wherein "Bz" represents a benzyl group, "Gal" represents a galactose residue, wherein "GlcNAc" represents an N-acetyl-D-glucosamine residue, and "β" represents an anomer assuming a cis configuration, of anomers of glycosidic linkage at position 1 of the sugar ring;
- (b) reaction pH: the protein has a high activity at or around neutral; and [[or]]

(c) divalent ion requirement: the activity is enhanced in the presence of at least Mn<sup>2+</sup> or Co<sup>2+</sup>.

## Claims 15-23 (canceled)

- 24. (withdrawn/currently amended) The method according to Claim 13, wherein the protein has at least one of the following properties (a) to (c):
- (a) acceptor substrate specificity: the protein transfers GlcNAc from UDP-GlcNAc to an acceptor substrate having an oligosaccharide residue in a quadruple-stranded form at the nonreducing end of an N-linked oligosaccharide through a β1,3 glycosidic linkage and synthesizes an oligosaccharide has a significant transferring activity for at least Bz-β-lactoside and/or Galβ1-4GlcNAc groups, wherein "Bz" represents a benzyl group, "Gal" represents a galactose residue, "GlcNAc" represents an N-acetyl-D-glucosamine residue, and "β" represents an anomer assuming a cis configuration, of anomers of glycosidic linkage at position 1 of the sugar ring;
- (b) reaction pH: the protein has a high activity at or around neutral; and [[or]]
- (c) divalent ion requirement: the activity is enhanced in the presence of at least Mn<sup>2+</sup> or Co<sup>2+</sup>.

## Claims 25-30 (canceled)

- 31. (new) The method according to Claim 13, wherein the protein comprises the amino acid sequence of SEQ ID NO: 2.
- 32. (new) The method according to Claim 13, wherein the protein comprises the amino acid sequence of SEQ ID NO: 16.
- 33. (new) The protein of Claim 14 comprising the amino acid sequence of SEQ ID NO: 2.

NARIMATSU et al. - Appln. No. 10/539,834

34. (new) The protein of Claim 14 comprising the amino acid sequence of SEQ ID NO: 16.